

*Def A1*

- 1 -

## SEQUENCE LISTING

<110> Instituto Científico y Tecnológico de Navarra (ICTN)

<120> TGF $\beta$ 1-inhibitor peptides

<160> 10

<210> SEQ ID NO: 1

<211> 15

<212> Peptide

<213> Artificial sequence

<220> Domain

<223> Derived from TGF $\beta$ 1, position 319-333

<400> His Ala Asn Phe Cys Leu Gly Pro Cys Pro Tyr Ile Trp

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Ser Leu

15

<210> SEQ ID NO: 2

<211> 14

<212> Peptide

<213> Artificial sequence

<220> Domain

<223> Derived from TGF $\beta$ 1, position 322-335

<400> Phe Cys Leu Gly Pro Cys Pro Tyr Ile Trp Ser Leu Asp

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Thr

<210> SEQ ID NO: 3

<211> 12

<212> Peptide

<213> Artificial sequence

<220> Domain

<223> Deduced as complementary to TGF $\beta$ 1, position 731-742

<400> Thr Ser Leu Asp Ala Thr Met Ile Trp Thr Met Met

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<210> SEQ ID NO: 4

<211> 15

<212> Peptide

<213> Artificial sequence

<220> Domain

<223> Overlapping with the extracellular region of the rat type III receptor, position 245-259

<400> Ser Asn Pro Tyr Ser Ala Phe Gln Val Asp Ile Ile Val

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Asp Ile

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<210> SEQ ID NO: 5

<211> 9

<212> Peptide

<213> Artificial sequence

<220> Domain

<223> Modification P54 deduced as complementary to TGF $\beta$ 1, position 731-742

<400> Thr Ser Leu Met Ile Trp Thr Met Met

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<210> SEQ ID NO: 6

<211> 14

<212> Peptide

<213> Artificial sequence

<220> Domain

<223> Derived from the modified human type III receptor, position 729-742

<400> Thr Ser Leu Asp Ala Ser Ile Ile Trp Ala Met Met Gln  
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Asn

<210> SEQ ID NO: 7  
<211> 14  
<212> Peptide  
<213> Artificial sequence  
<220> Domain  
<223> Derived from the modified human type III receptor,  
position 241-254  
<400> Ser Asn Pro Tyr Ser Ala Phe Gln Val Asp Ile Thr Ile  
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Asp

<210> SEQ ID NO: 8  
<211> 15  
<212> Peptide  
<213> Artificial sequence  
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<223> Position 247-261 of endoglin  
<400> Glu Ala Val Leu Ile Leu Gln Gly Pro Pro Tyr Val Ser  
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Trp Leu  
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<210> SEQ ID NO: 9  
<211> 15  
<212> Peptide  
<213> Artificial sequence  
<220> Domain  
<223> Position 445-459 of endoglin

<400> Leu Asp Ser Leu Ser Phe Gln Leu Gly Leu Tyr Leu Ser  
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Pro His

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<210> SEQ ID NO: 10

<211> 23

<212> Peptide

<213> Artificial sequence

<220> Domain

<223> Modification P12, position 322-335 of TGF $\beta$ 1

<400> His Glu Pro Lys Gly Tyr His Ala Asn Phe Cys Leu Gly

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Pro Cys Pro Tyr Ile Trp Ser Leu Asp Thr

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20